

Fig. 1

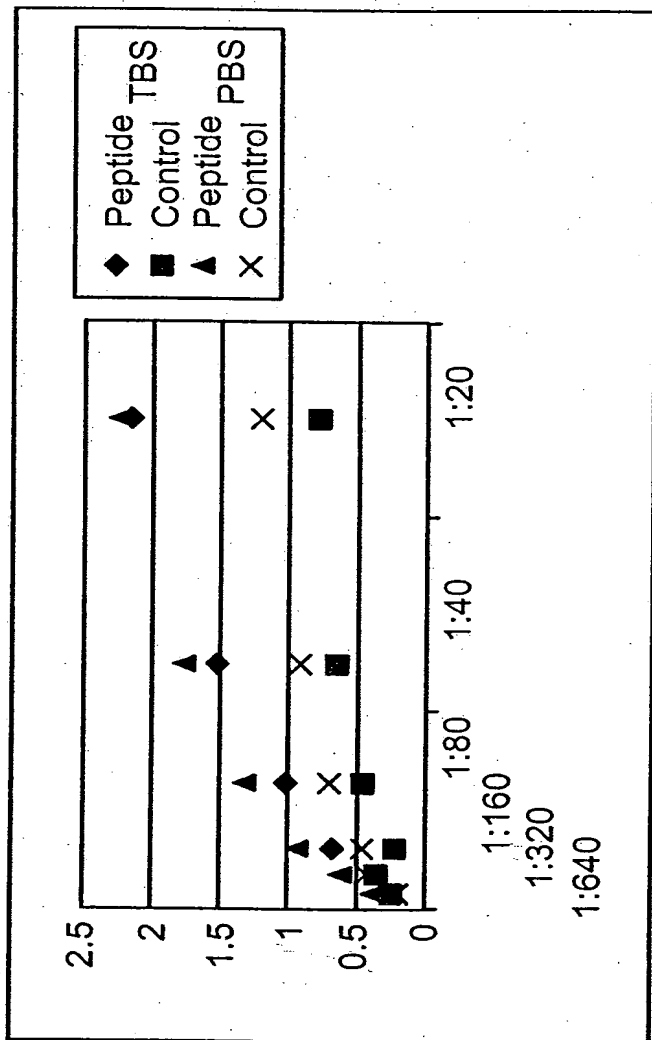


Fig. 2

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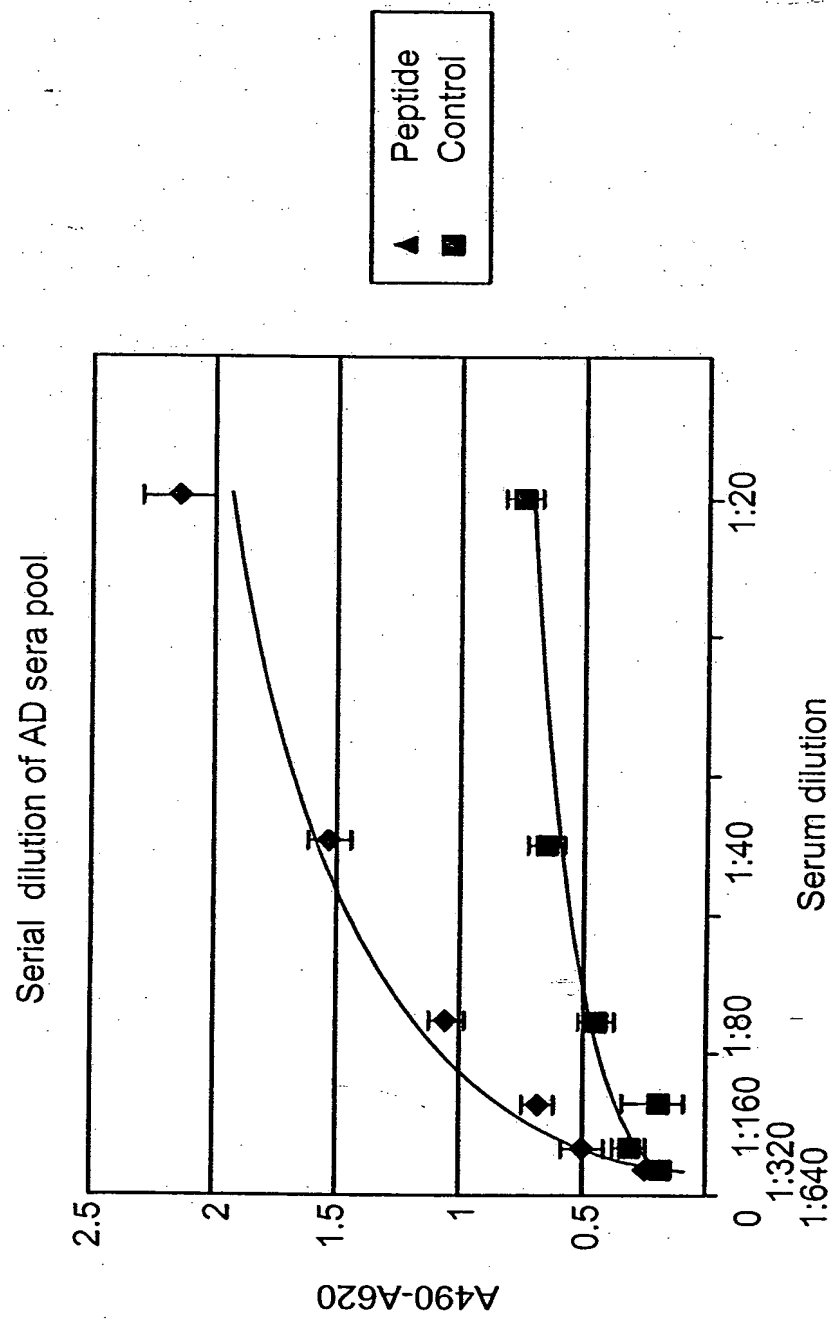


Fig. 3

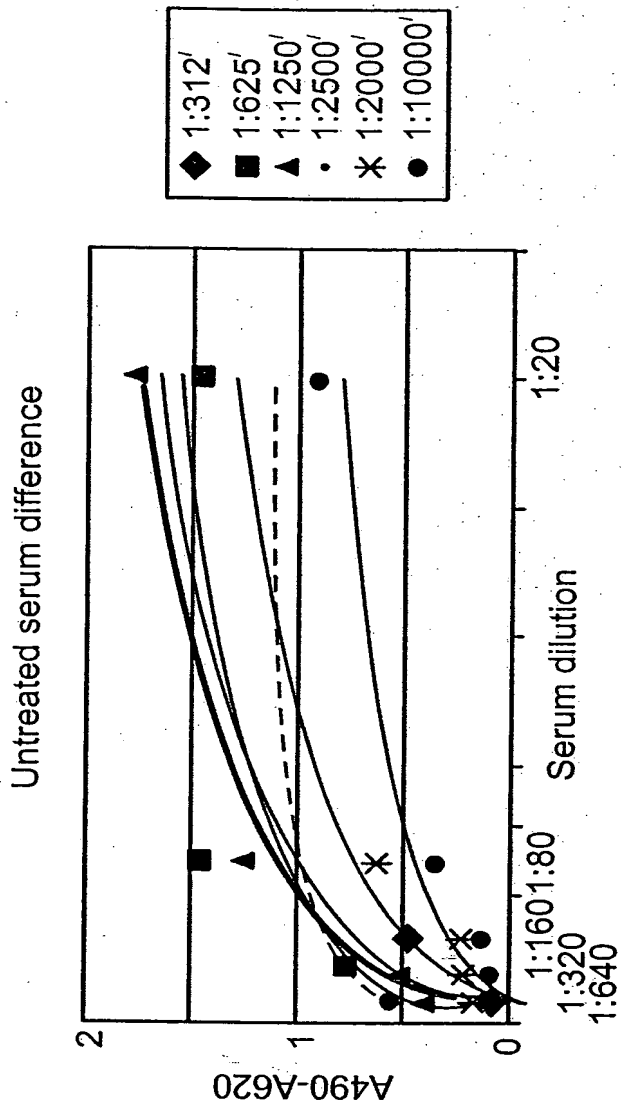


Fig. 4

Met Ala Glu Pro Arg Gln Glu Phe Glu Val Met Glu Asp His Ala  
5 10 15  
Gly Thr Tyr Gly Leu Gly Asp Arg Lys Asp Gln Gly Gly Tyr Thr  
20 25 30  
Met His Gln Asp Gln Glu Gly Asp Thr Asp Ala Gly Leu Lys Glu  
35 40 45  
Ser Pro Leu Gln Thr Pro Thr Glu Asp Gly Ser Glu Glu Pro Gly  
50 55 60  
Ser Glu Thr Ser Asp Ala Lys Ser Thr Pro Thr Ala Glu Asp Val  
65 70 75  
Thr Ala Pro Leu Val Asp Glu Gly Ala Pro Gly Lys Gln Ala Ala  
80 85 90  
Ala Gln Pro His Thr Glu Ile Pro Glu Gly Thr Thr Ala Glu Glu  
95 100 105  
Ala Gly Ile Gly Asp Thr Pro Ser Leu Glu Asp Glu Ala Ala Gly  
110 115 120  
His Val Thr Gln Ala Arg Met Val Ser Lys Ser Lys Asp Gly Thr  
125 130 135  
Gly Ser Asp Asp Lys Lys Ala Lys Gly Ala Asp Gly Lys Thr Lys  
140 145 150  
Ile Ala Thr Pro Arg Gly Ala Ala Pro Pro Gly Gln Lys Gly Gln  
155 160 165  
Ala Asn Ala Thr Arg Ile Pro Ala Lys Thr Pro Pro Ala Pro Lys  
170 175 180  
Thr Pro Pro Ser Ser Gly Glu Pro Pro Lys Ser Gly Asp Arg Ser  
185 190 195  
Gly Tyr Ser Ser Pro Gly Ser Pro Gly Thr Pro Gly Ser Arg Ser  
200 205 210  
Arg Thr Pro Ser Leu Pro Thr Pro Pro Thr Arg Glu Pro Lys Lys  
215 220 225  
Val Ala Val Val Arg Thr Pro Pro Lys Ser Pro Ser Ser Ala Lys  
230 235 240  
Ser Arg Leu Gln Thr Ala Pro Val Pro Met Pro Asp Leu Lys Asn  
245 250 255  
Val Lys Ser Lys Ile Gly Ser Thr Glu Asn Leu Lys His Gln Pro  
260 265 270  
Gly Gly Gly Lys Val Gln Ile Ile Asn Lys Lys Leu Asp Leu Ser  
275 280 285  
Asn Val Gln Ser Lys Cys Gly Ser Lys Asp Asn Ile Lys His Val  
290 295 300  
Pro Gly Gly Gly Ser Val Gln Ile Val Tyr Lys Pro Val Asp Leu  
305 310 315  
Ser Lys Val Thr Ser Lys Cys Gly Ser Leu Gly Asn Ile His His  
320 325 330  
Lys Pro Gly Gly Gly Gln Val Glu Val Lys Ser Glu Lys Leu Asp  
335 340 345  
Phe Lys Asp Arg Val Gln Ser Lys Ile Gly Ser Leu Asp Asn Ile  
350 355 360  
Thr His Val Pro Gly Gly Gly Asn Lys Lys Ile Glu Thr His Lys  
365 370 375  
Leu Thr Phe Arg Glu Asn Ala Lys Ala Lys Thr Asp His Gly Ala  
380 385 390  
Glu Ile Val Tyr Lys Ser Pro Val Val Ser Gly Asp Thr Ser Pro  
395 400 405  
Arg His Leu Ser Asn Val Ser Ser Thr Gly Ser Ile Asp Met Val  
410 415 420  
Asp Ser Pro Gln Leu Ala Thr Leu Ala Asp Glu Val Ser Ala Ser  
425 430 435  
Leu Ala Lys Gln Gly Leu (SEQ ID NO: 71)  
440

Fig. 5

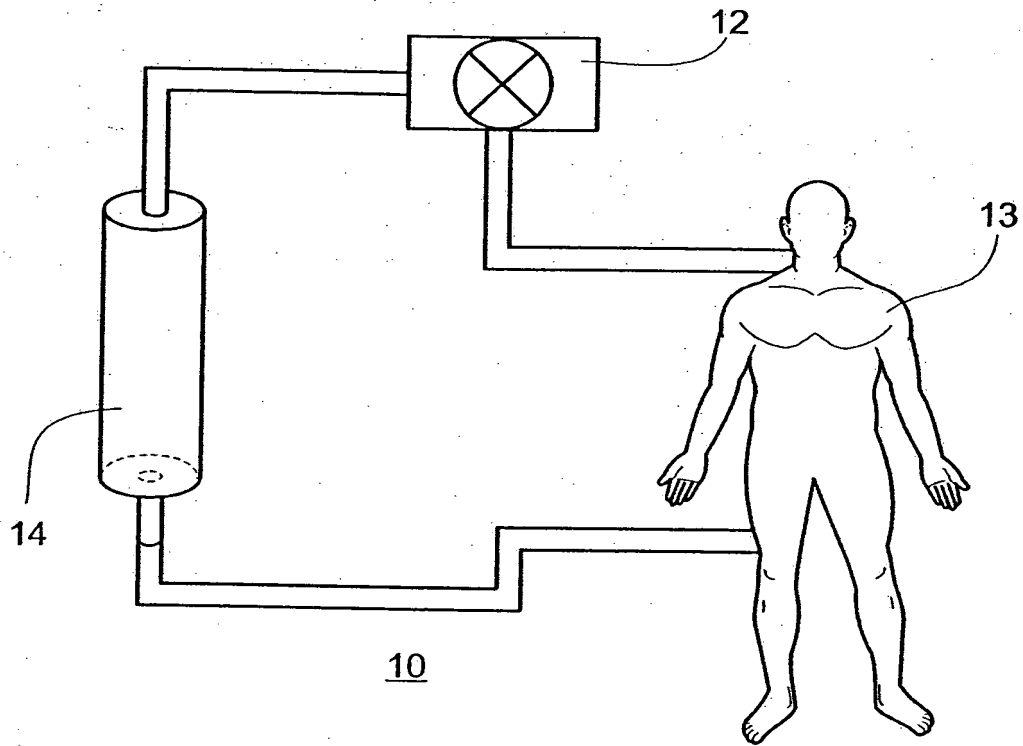


Fig. 6

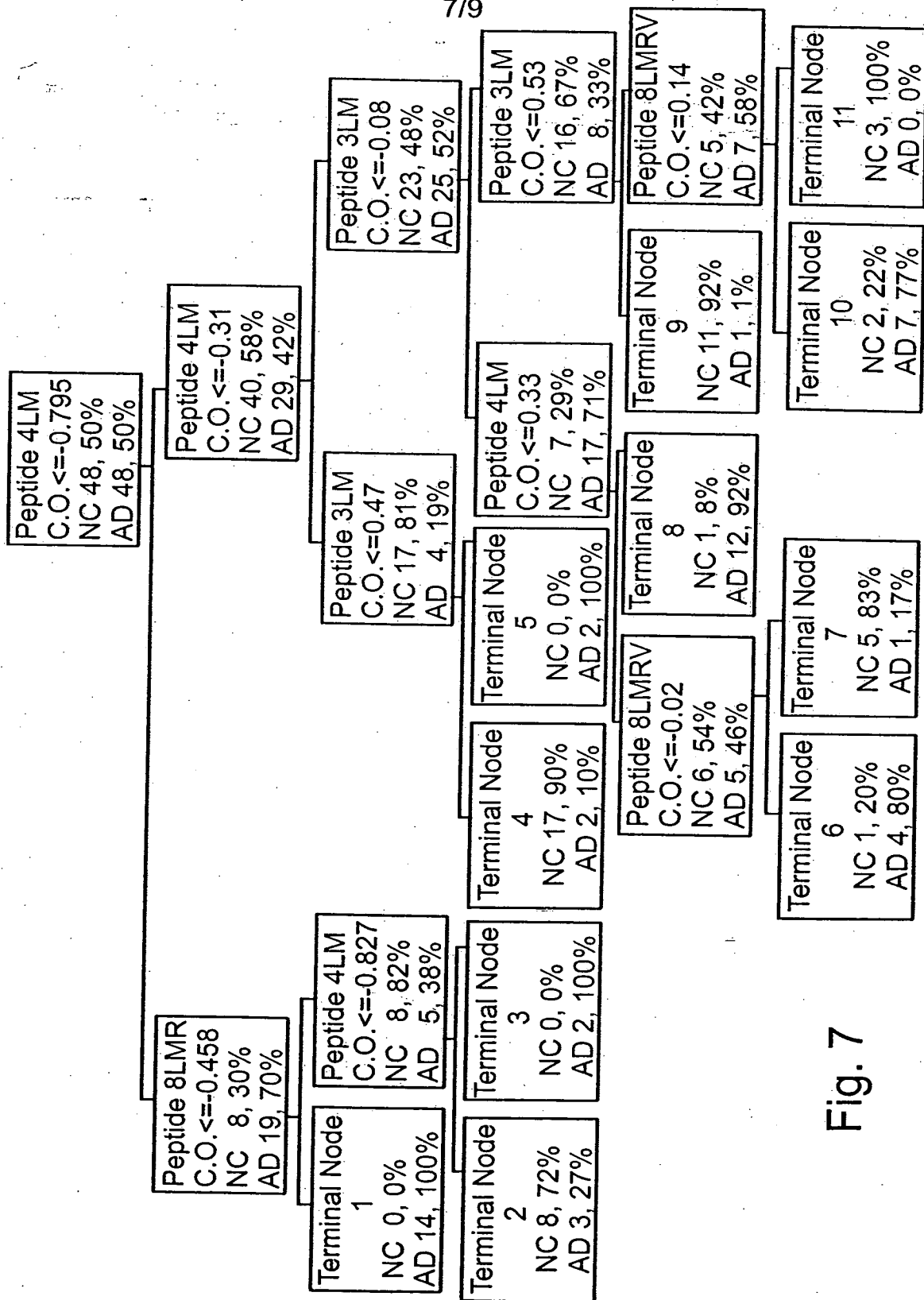


Fig. 7

Antibody profiles characteristic for AD or NC sera

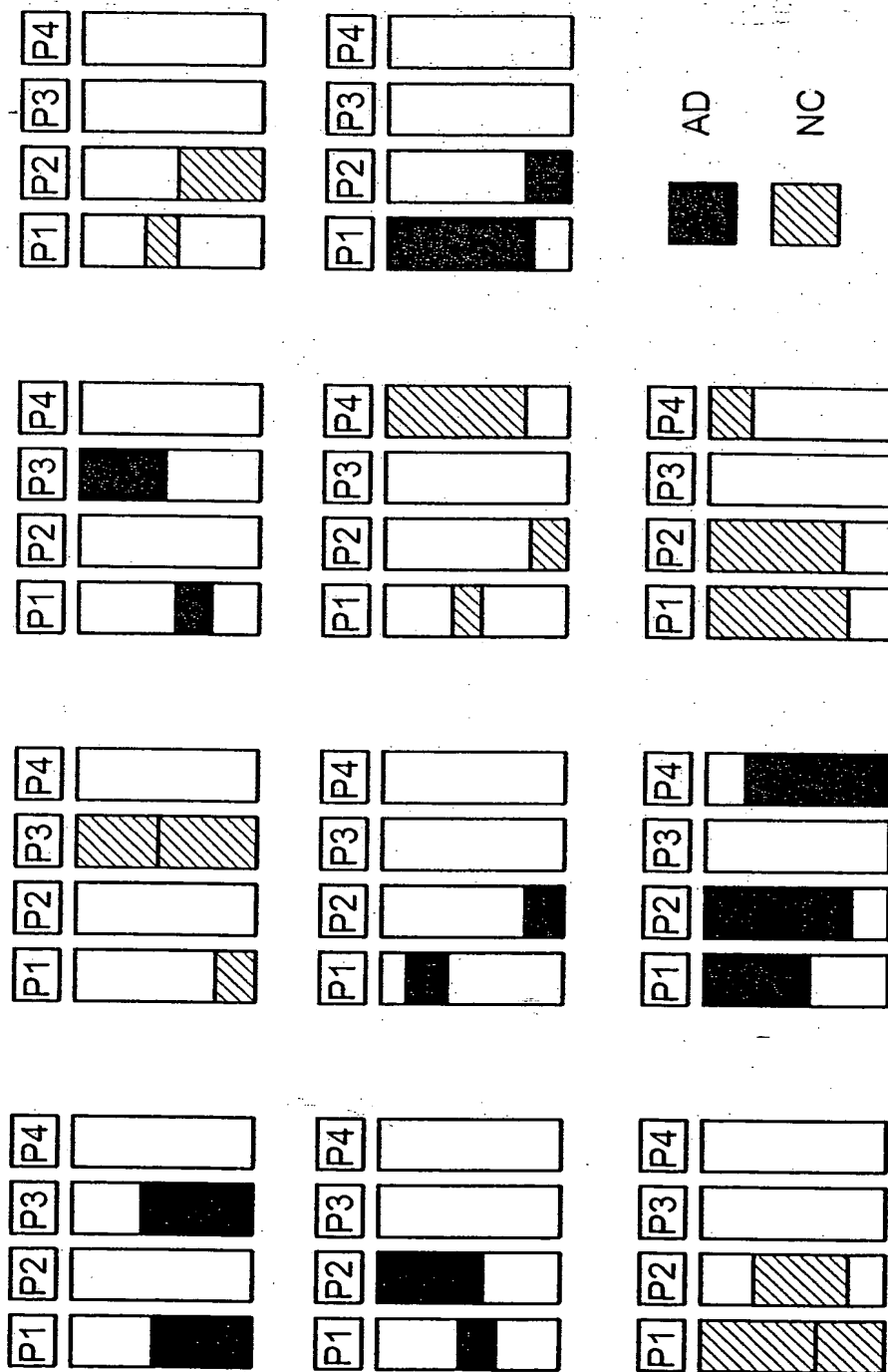


Fig. 8

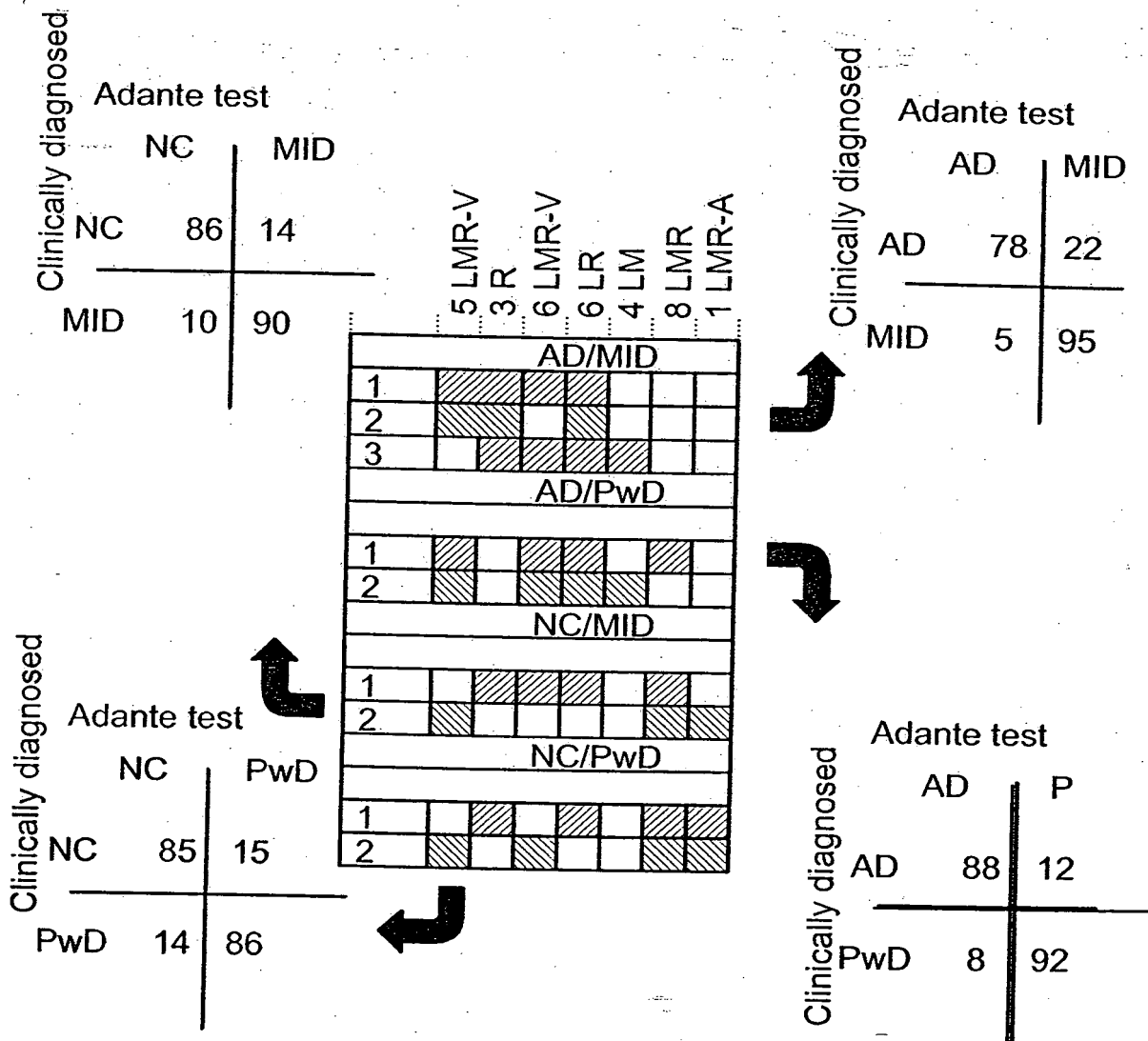


Fig. 9